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SEO 31



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SEO 34



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SEQ 43



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SEO 49



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<213> Human immunodeficiency virus type 1



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<213> Human immunodeficiency virus type 1



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SEO 78

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SEO 91



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<213> Human immunodeficiency virus type 1

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ggatgtaatc agtttatggg atcaaagcct aaagccatgt gtaaagttga ccccactctg 780
tgtcacttta aactgtagta aggttaccaa taatgctact tacaataata ctgatgatat 840
aaaaaattgc tcttttaatg caaccacaga aataagagac aagaaacgca aagagtatgc 900
actgttttat agactcgata tagtaccact aaatgagaat aagaacagct ctagtaacta 960
tagtgagtac atattaataa attgtaatac ctcaaccata acacaagcct gtccaaaggt 1020
```

SEO 94



```
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<211> 2651
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gattaggagt aggagcatta atagtagcat taatcttagc aataattgtg tggaccatag 180
catatctaga atataggaaa ttgttaagac aaagaaaaat aaacaggtta attgaaagaa 240
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gcaaccacag aaataagaga caagaaacgc aaagagtatg cactgtttta tagactcgat 780
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                                     SEO 95
```



```
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qtatgtacaa gaccaacaa taacacaaga aaaagtataa ggatagggcc aggacaagca 1200
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aattgtggag gagaattett ttattgtaat acatcaagae tgtttaaegg tacatacaat 1440
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gcagcgagag cggtggaact tctgggacgc agcagtctca ggggactaca gagggggtgg 2640
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<210> 106
<211> 261
<212> DNA
<213> Human immunodeficiency virus type 1
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<400> 106
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caaagaaaaa taaacaggtt aattgaaaga attagggaaa gagtagaaga cagtggcaat 180
gagagtgagg gggatactga ggaattgtca acactggtgg atatggggaa tcttaggctt 240
ttggatgcta atgatttata a
<210> 107
<211> 2574
<212> DNA
<213> Human immunodeficiency virus type 1
<220>
<223> isolate=94IN476.104; gene=env
<400> 107
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                                    SEO 96
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ttttggatgc taatgattta taatgtagta qqqaacttgt qqqtcacagt ctattatggg 120
gtacctgtgt ggaaagaagc aaaaactact ttattctgtg catcagatgc taaagcttat 180
gagaaggagg tgcataatat ctgggctaca catgcctgtg tacccacaga ccccaaccca 240
caagagatgg atttagtaaa tgtaacagaa aattttaaca tgtggaaaaa tgacatggtg 300
gatcagatgc atgaggatgt aatcagttta tgggatcaaa gcctaaagcc atgtgtaaag 360
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cgcaaagagt atgcactgtt ttatagactc gatatagtac cactaaatga gaataagaac 540
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<211> 624
<212> DNA
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<220

<223> isolate=94IN476.104; gene=nef

<400> 108

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<213> Human immunodeficiency virus type 1

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acttataaag gagcattcga tctcqccttc tttttaaaag aaaaqqqqqq actqqatggg 300
ttaatttact ctaagaaaag gcatgaaatc cttgatttat gggtttataa cacacaaggc 360
tacttccctg attggcaaaa ctacacacca ggaccagggg tcagatatcc actgaccttt 420
ggatggtgct acaagctagt accagttgac ccaagtgtag tagaagaggc caacaaagga 480
gaaaacaact gtttgctaca ccctatgagc caacatggaa tggatgatga agatggagaa 540
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ccggagtatt acaaagactg ctga
<210> 109
<211> 17
<212> DNA
<213> artificial sequence
<220>
<223> Upstream primer UP1A spanning tRNA primer binding state
<400> 109
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                                                                   17
<210> 110
<211> 22
<212> DNA
<213> artificial sequence
<220>
<223> Downstream primer Low2 spanning the R/U5 junction in the
      3' long terminal repeat
<400> 110
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<210> 111
<211> 28
<212> DNA
<213> artificial sequence
<220>
<223> Upstream UP1AMlu1 primer containing Mlu1 restriction site
<400> 111
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                                                                   28
<210> 112
<211> 35
<212> DNA
<213> artificial sequence
<220>
<223> Downstream LowlMlul primer containing Mlul restriction site
<400> 112
accagacgcg tacaacagac gggcacacac tactt
                                                                   35
                                     SEO '98
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